

AGATCTACACAAGCAAATTGAAAAATAGATAAAATTTCGAGGTATTAAAGCCGACTAAACAAATGAGTGAAGAA 80
 TCTAGATGTGTTCCGTTAACCTTTTATCTATTTAAAAGCGTCCATAATTCGGCTGAATTGTTACTCACTTCTT

 GAAAAGAAAAAAATAAAATACATATTTGAGTTAGTAAAAGAGAAAGAAAAATAAGAAGACCTCGGCTAACAGTCGA 160
 CTTTCTTTTATTATGTATAAAACTCAATCATTCTCTTCTTTTATTCTGGAGCCGAATTGTCAGCT

 AAAACCAGAAATAATAAAAGAAAGAGACTGTGATTTTAATGGAAATCGTGAGGAAAAGAAAATTAACTTTCAATT 240
 TTTGGTCTTATTATTTCTGACACTAAAATTACCTTAGCACTCCTTCTTTAAAATTAAAGTAAAA

 CGAGGGATTAATTGTTGAAGTTGATGAAAAATCTAGATAAAAATGCAGATCAAAATGTGTTGAATTGACATTATT 320
 GCTCCCTAATTAAACACATTCAACTACTTTAGATCTATTTTACGTCTAGTTTACACAACTTAACTGTAATAA

 GAAATACGTAGTATATCAATAATGGGGTTGCTATTTATTTGCGAAGATTGAAAATCTGAGTGAAGAAAATAGTT 400
 CTTTATGCATCATATAGTTATTACCCCCAACAGATAAAATAACGCTTCTAACCTTAGACTCACTTCTTTATCAA

 TGCGAGAGCAAAAAACCCCTGCCGTTTTCAAATGACTTGGAAAAATTCAATTGTGAGCGGTAGCGAAACTTGAA 480
 ACGCTCTCGTTTTGGAACGGCAAAAAAGTTACTGAAACCTTTTAAGTAACACTGCCATCGCTTGAAACTT

 ATTTTTACATTGAAATTGAAAAATAAGGCAAAAGAACTCAAATGGAAAAATTATTTATAAAAAAGGAGATCG 560
 TAAAAAAATGTAACCTTAAACTTTTATTCCGTTTCTTGAGTTACCTTTTATAATAATTTTCTCTAGC

 Asn Phe Leu His Trp Phe Glu Lys Ile Arg Gln Lys Leu Lys Trp Lys Lys Tyr Tyr Lys Arg Arg Ser

 GATATGGATTTAAAGCAGAAACTGACATTGAATGAAAAAGATTGGAAAAATCTATGCTGAGAGTGAATTAAA 640
 CTATAACCTAAATTTCGTCTTTGACTGTAACCTACTTTTCTAAACCTTTAGATAACGACTCTCACTTAATT

 Asp Met Asp Phe Lys Ser Arg Lys Leu Thr Leu Asn Glu Lys Lys Asp Leu Glu Lys Ile Tyr Ala Glu Ser Glu Leu Lys

 AGCAAAAAATTGGGAACTCAACCCGGTGTGTTAGAAATGACGATGAAAGAAATGATGAAAATATCAACCTCGATG 720
 TCGTTTTTAACCTTGAGTTGGGCCACAACAAATCTTACTGCTACTTCTTACTACTTTATAGTTGGAGCTAC

 Ala Lys Lys Leu Gly Thr Gln Pro Gly Val Val Leu Glu Met Thr Met Lys Asn Ile Asn Leu Asp

 TTAATGAAGAAACAGCAGGTCAATATAGGAAATTCAAAAATAAGTTGAGCATAGTAAATCAGATGATCTAGTAACG 800
 AATTACTTCTTGTGTCAGTTATATCCTTAAATAAGTTTATTCAACTCGTATCATTAGTCTACTAGATCATGCC

 Val Asn Glu Glu Thr Ala Gly Gln Tyr Arg Lys Leu Phe Lys Asn Lys Val Glu His Ser Lys Ser Asp Asp Leu Val Thr

FIGURE 1A

GGACTATTAGAGTGTGGAACTCGAAATAGTTTGATAAAACAAGAAGTGCCTTCGTTTGTATTGTGAGAGAATTCA 880
 CCTGATAATCTCACACCTTGAGCTTATCAAACATTTGTTCTCACGGAAAGCAAAACATAAACACTCTCTTAAGT
 Gly Leu Leu Glu Cys Gly Thr Arg Asn Ser Phe Asp Lys Thr Arg Ser Ala Phe Arg Phe Cys Ile Cys Glu Arg Ile Gln
 GCAACTGAGAAAAGAAGCTGATAATGCAAGAAGAGTAAAGATTCGATACAATGAAAGCAAAACTAAAGAGGTTTG 960
 CGTTGACTCTTCGACTATTACGTTCTCTCATTCTAAAGCTATGTTACTTCGTTTGATTCTCCGAAAC
 Gln Leu Arg Lys Glu Ala Asp Asn Ala Arg Arg Val Lys Asp Phe Asp Thr Met Lys Ala Lys Thr Lys Glu Ala Phe
 AATTGAGTTTGTTTGATAAGGATTTTGAGTGAAATAGAATTCAATGGAATGATATTCTCACACACAAAAAGAC 1040
 TTAACTCAAACAAACACTATTCTAAACACTCACTTTATCTTAAGTTACCTACTATAAGAGTGTGTTCTG
 Glu Leu Ser Phe Val Phe Asp Lys Asp Phe Leu Ser Glu Asn Arg Ile Gln Trp Asn Asp Ile Ser His Asn Lys Lys Asp
 TCTGCAAGTAAAGAAAAACATGAAAGAAGCGGACACAATGGATGATTTTAAGAGGCTAAAAAATAATAATCTAC 1120
 AGACGTTCATTTCTTTGTTACTTCTCGCTGTGTTACCTACTATAAAATTCTCGATTTTTATTATTAGATG
 Ser Ala Ser Lys Arg Lys Thr Met Lys Glu Ala Asp Thr Met Asp Asp Ile Phe Lys Arg Leu Lys Asn Asn Lys Ser Thr
 ATATGATCGTTATGCTGGATTCTTCTATTGTTCGATTACAGGTTGCAGACCAGCAGAAGTTAAAGGGTATAGAGA 1200
 TATACTAGCAATACGACCTAAGGAAAGATAAACAGCTAATGTCCAACGTCTGGTCGTTCAAATTTCCATATCTCT
 Tyr Asp Arg Tyr Ala Gly Phe Leu Ser Ile Cys Ser Ile Thr Gly Cys Arg Pro Ala Glu Val Leu Lys Gly Ile Glu
 TAGTAAGAAACAGATATGAGGATGGTATATCTTTAAAATCTTGGTGCAAAGGTTGAAATGACAGAGGGCAAAGCGAA 1280
 ATCATTCTTGCTATACTCCTACCATATAGAAAATTGAAACCACGTTCCAACCTTACTGTCTCCGTTCGCTT
 Ile Val Arg Asn Arg Tyr Glu Asp Gly Ile Ser Phe Lys Ile Leu Gly Ala Lys Val Gly Asn Asp Arg Gly Gln Ser Glu
 AGAACATTACATTTGATTATCAAATATCATGATAATGAGCAAATGAATTATTTGTCGCAATTAAAGATAATAA 1360
 TCTTGTAAATGAAACTAAATAGTTATAGTACTATTACTCGTTACTTAATATAAACAGCGTTAATTCTATTATT
 Arg Thr Leu His Phe Asp Leu Ser Lys Tyr His Asp Asn Glu Gln Met Asn Tyr Ile Leu Ser Gln Leu Lys Asp Asn Lys
 ATTTTCTACAAACCAGATGGGAAGCTCTACAAACAGCTTGAGGAATACCTCTACATCCAACATAGAACGTTTCACTGT 1440
 TAAAAAGATGTTGGTCTACCCCTCGAGATGTTGCGAACCTCGTTATGGAGATGTAGGTTATCTTGCAAAAGTGACA
 Phe Phe Tyr Lys Pro Asp Gly Lys Leu Tyr Asn Ser Leu Arg Gln Tyr Leu Tyr Ile Gln His Arg Thr Phe Ser Leu
 ATACACTTCGTACAGGGTTGCGAGTGATCTCAAGGCATCCGGTGCAGATGACTTCACCATAGCGGCTNTTTGGTCAC 1520
 TATGTGAAGCAGTGTCCCAACGCTACTAGAGTCCGTAGGCCACGTCTACTGAAGTGGTATCGCCGAAAACCCAGTG
 Tyr Thr Leu Arg His Arg Val Ala Ser Asp Leu Lys Ala Ser Gly Ala Asp Asp Phe Thr Ile Ala Ala ??? Leu Gly His

FIGURE 1B

AGAGTGACTCAAAGCCAGGAGTTACTACGGCTATGCTCGTCGNAAGGTGGTATCGCTGTA
TCTCACTGAGTTCGGT CCTCAATGATGCCGATACGAGCAAGCAGCNTTCCACCA1AGCGACATTGACCACAACTCACGA 1600

Arg Val Thr Gln Ser Gln Glu Leu Leu Arg Leu Cys Ser Phe Val ??? Arg Trp Tyr Arg Cys Asn Trp Cys

CTGATGTTGTGAAAGCAAACAAGAGTCAGTTNGCTGTATCAAGGACTCCGAGCCAGATCT 1660
GACTACAACACTTTCGTTGTTCTAGTCAGTCAANGACATAGTTCCCTGAGGCTCGGTCTAGA

FIGURE 1C

AGATCTAACCAAGTTAAAATCGCACTTCAAGAAGTAAAATAGGGCCGGCACCGGCTCTTTTTGGTGTGGTAG
 80
 TCTAGAGTTGGTCAAATTTAGCGTGAAGTCTTCATTTTATCCCCGGCGAGAAAAAAACACAAAAACATC

TTAGTGGATATCTGTTAGCTACAGAGAAAAGCGATTTAGAGGGTTGACGAGGTTTCGAGCTATCCAGGGTTT
 160
 AATCACCTATATAGACAATCGATGTCTTTCGCTAAAATCTCCAAACTGCTCCAAAAAGCTCGATAGGTCCAAAA

TGGGTTTTGGGTTGGATCAGAAAAGTCGTTCAAGATTATTGACATAAAGACAGGAAGGTTATAACAAGTACCAAGATA
 240
 ACCCAAAAACCCAACCTAGTCTTCAGCAAGTCTAATAACTGTATTCTGCCTTCAAATATTGTTCATGGTCTAT

CGACAAAACAGCTTGCAAGGCTGGCTTGAAGGACTAAAAGAAGTGGGACTTCTTGAGTCTTGAATCAAGTTGGTC
 320
 GCTGTTTGGTCGAAACGTCCGACCGAAACTCCTGATTTCTCACCCCTGAAGAAACTCAGAACATTAGTTCAACCAG

AGAAACTCGATTACGATTGTAAGTAGAAATCTAACTCACATTGCAAGAAAGTCAAACCTACCTCTTAGTTACAACCAA
 400
 TCTTGAGCTAATGCTAACATTCTAGTTAGATTGAGTGAAAGCGTCTTCAGTTGAATGGAGAATCAATGTTGAGT

AAATTCCTAGCCTTTCAAGATCCTTAAGCATACATATTGTTAAACCGATTGTGTCCGGTGTGGAGCCAT
 480
 TTTAAAGGATCGGAAAGTCTAGGAATTGATGTATAAAACAAATTGGCTAACACAGGCCACAAACACACCTCGGTA

TGATCCGAGTGGTCAATATGTGATTGTCGCCAACAGTGTATGTAGGTCTAAACGGGAGTGCTACAAAAGACCATACC
 560
 ACTAGGCTCACCAAGTTACACTAACAGCGGTTGTCACATACATCCAGATTGCCCTCACGATGTTCTGGTATGG

CGAAACGAGTGCCTAAGTGTGGTTATCAACCAGGTAAAGCTATGAGAAAGCCCAGCCATAATGGGGTAGGTTGAAG
 640
 GCTTGCTCACGGATTCAAAAACCAATAGTTGGTCCATTGATACTCTTCGGGTCGGTATTACCCCAATCCAACCTC

CAAGTCTTCATATGGTGCACACAAGGGGTGTAGTAGGGTGTGTCGTCAAACTGAAAGGTTGATAGCTCAAGCTTGCT
 720
 GTTCAGAAGTATACCACGCTGTGTTCCCCACATCATCCCACAGCAGTTGACTTCCAAACTATCGAGATTGAAACACGA

TCTGTGGGTCAAGCCTCAAGTGTGATCTGTTGTCGTCACCTGATAACTTCACTTTGAGTGAAATTCAAGGAGG
 800
 AGACACCCAGTTGGAGTTACGACTAGACACCACAGCAGATGGACTATTGAAAGTGAAAAGCTCACTTAAGTCCTCC

CGAAACTATGGGTCAAGCCCAGCTTGCTGGGGTCGGCACATCCAGCTTACAGCATTGGTGTCTTGCGAAGCTGAAGC
 880
 GCTTGATACCCAGTTGGGTGAAACGACCCCAAGCCGTGTAGGTGAAATGTCGTAACCACGAGAACGCTTCGACTTCG

FIGURE 1D

ACAAAAATCTAATCCAGGGTTGGGTTTTATACCAAGCAAAACAAAAAATAAAACAAGAAAATTCGAGCGA 960
TGTTTTAGATTAGGTCCCAACCCAAAAATATGGTCTCGTTGTTTTATTTGTTCTTTAAAAGCTCGCT

AAAAATATTTGGAATTTAAAGGCATACTGCTACCGCACTTTGCCATATTAAAACCTGACTATCTTATAAGT 1040
TTTTATAAAACCTTAAAAAATTCCGCTATGAACGATGGCGTGAAACGGTATAAATTGGACTGATAGAAATTCA

TAATAGATATATCCGTTAGATTATAAGTATGTTAAAACGAGTAAAAACAATAACTTATATTTAATTCTGAATTATA 1120
ATTATCTATATAGGCAATCTAATATTCACTACAATTGCTCATTTGTTATTGAATATATAAATTAAAGACTTAATAT

TTTGACAGTGATTATTAATATTAAGAGATATCTATTAGCTAAATATAACTAAAAAGAGGTAATATGGAT 1200
AAACTGTCACTAATAAATTATATAATTCTCTATAGATAATCGAATTATGATTTCTCCATTATACCTA

TGTGTATTTAAAAAGCATTAGAAAATGAAATAGAACATTATAAAAAGACGGTATATCAAATCTTCTAACATACTT 1280
ACACATAAATTTCGTAATCTTACTTATCTGTAATTGTTCTGCCACTATAGTTAGAAAGAATGTTATGAA

GCATTACTTGATATAGATAAAGCATTAAATGGTGATGAATGTGGCGATATTATAACTCAAATTATCATTGATGAAA 1360
CGTAATGAAACTATATCTATTTCGTAATTACCACTACTTACACCGCTATAATATTGAGTTAAAGGTAACTACCTT

GTTTGATCTTCTTGATGTTGAGCACAATTTCGGCTGGGCTTCAATAAAATAACAGAGACGAAATGAATATTCA 1440
CAAAACTAGAAGAACTACAACCTCGTAAAGCCGACCCGAAAGTTATTATGTCTGCTTACTTATAAATAGT

TCAGCTAAAATGAAAATGATTTAAAAAAACTCGTTCTTATTCACTCGATCAATTGGAGAAATTAAATTACGATGA 1520
AGTCGATTTGACTTTACTAAAATTGAGCAAGAAATAAGTAAGCTAGTTAACCTCTTAAATTATGCTACT

GATGAGTACAATACATCAAGAAATGATTAAGGATTAGATAATTACACATATGGAGAAATAACCATATGAATAATAAAT 1600
CTACTCATGTTATGTTAGCTTACTAATTCTTAATCTATTGTATACCTCTTATTGGTATACTTATTATTTA

AAGAGAATATATTGATTTGAAATAACAAAGATATAAAAGAAAGTCAGCTCTTAAAATATCGCATTGATCGATGTTT 1680
TTCTCTTATATAACTAAAGCTTATTGTTCTATATTCTTCACTGAGAATTGAGACGTAACAGCTACAAA

TAAAAGTAGATGAAAATTATTGATGAAGAGGATTGCAACTAAAGATATTGAAAATATCGTATGAAAATCCTATTGAT 1760
ATTTTCATCTACTTTAAATAACTACTTCTCTAACGTTGATTCTATAACTTTATAGCATACTTTAGGATAACTA

GATCCAGATGATGGCATAAGAAATCACAATTGCAACGAAGAAATGCCATGCTTCCGCTTAAAGCAAGCAAAA 1840
CTAGGTCTACTACCGTATTCTTTAGTGTAAAGCGTGCTCTTACGGATACGAAAGGCGTAATTGTTGTTCTT

FIGURE 1E

GAGATCT
←→ 1847
CTCTAGA

FIGURE 1F

Growth of ps^- vs ps^+ in ML5 at 28C, 32C & 36C

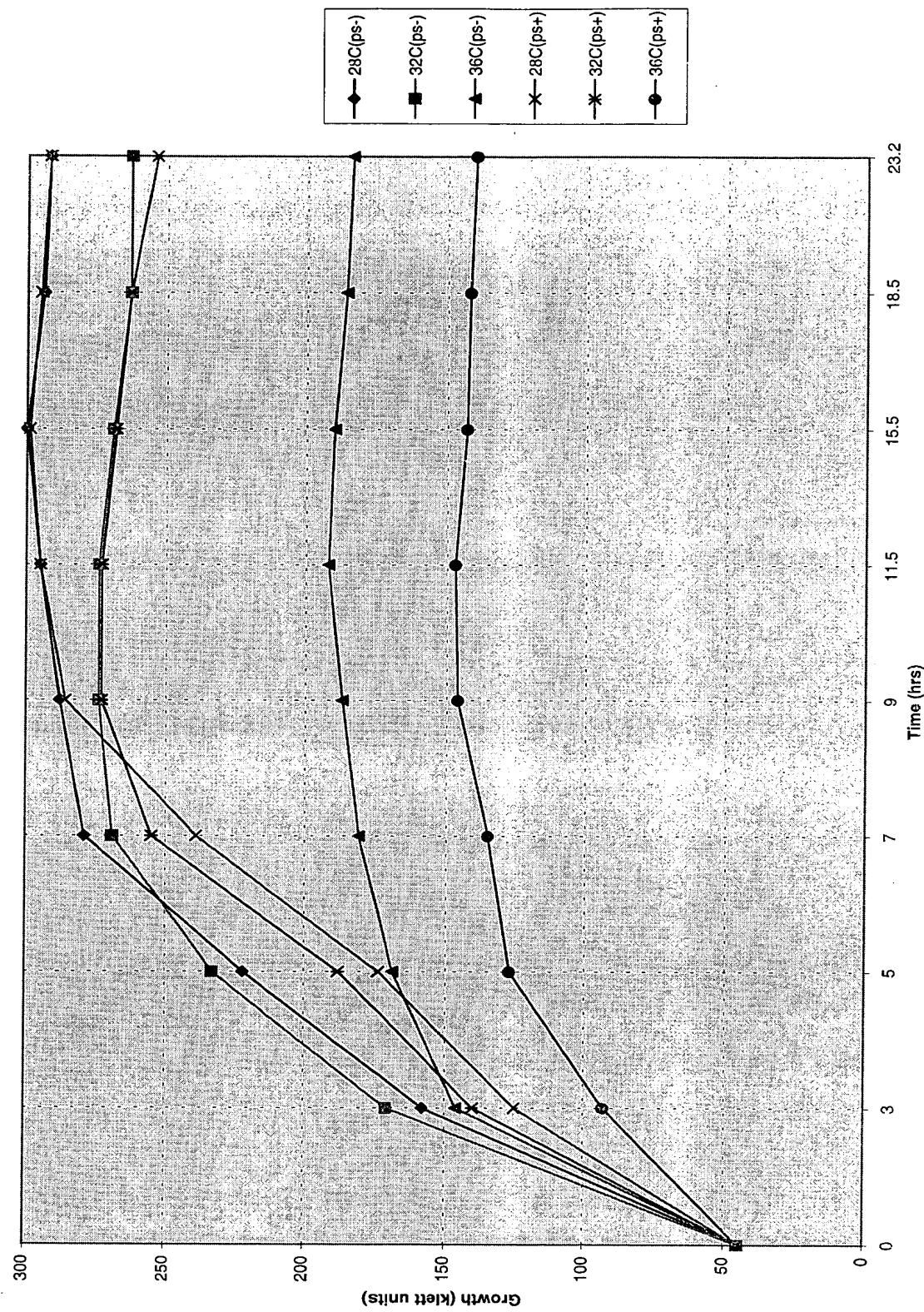


FIGURE 2

Growth of $ps(-)$ vs $ps(+)$ in LB at 28C,32C & 36C

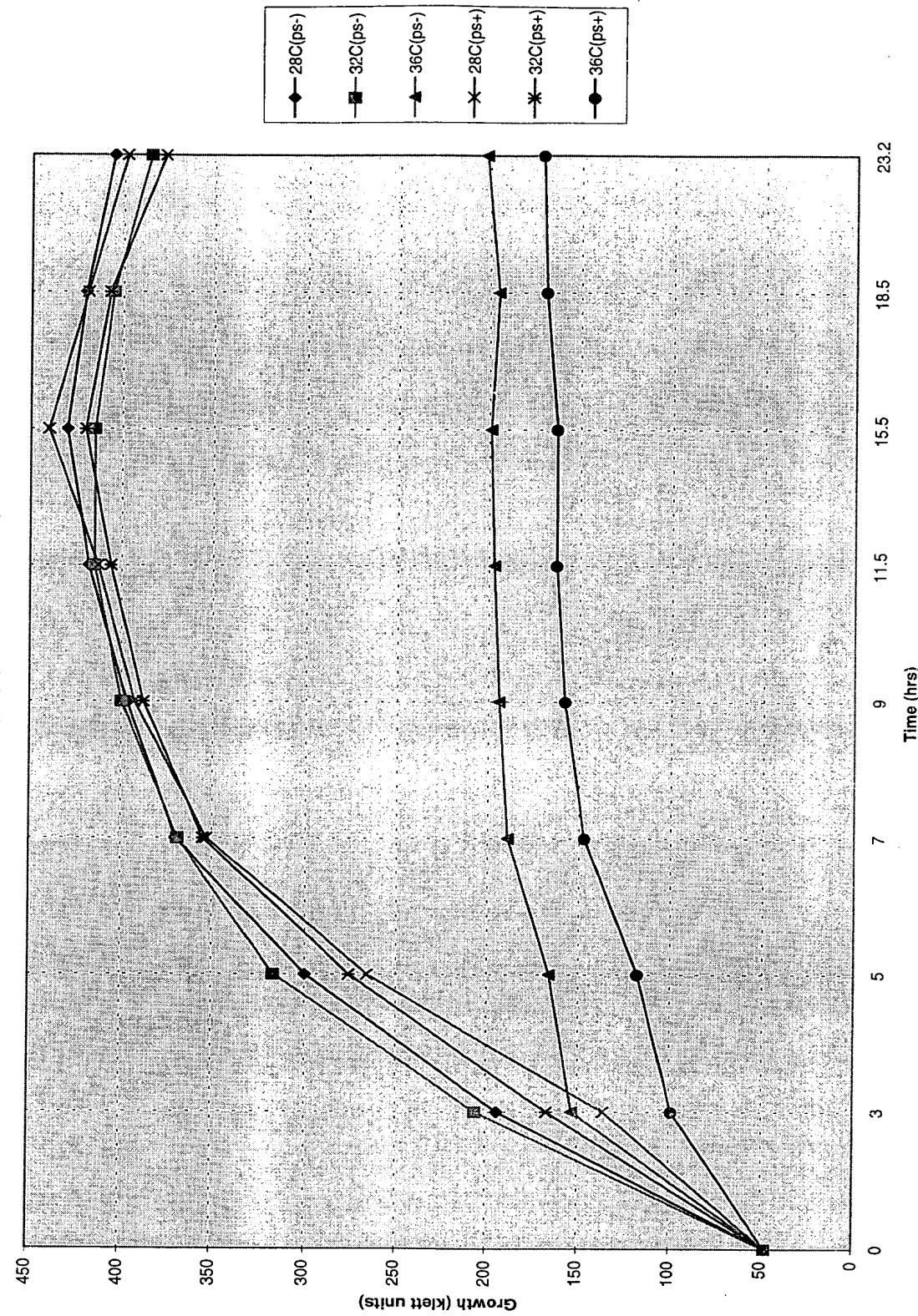


FIGURE 3